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Sequence Listing was accepted.

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Reviewer: Anne Corrigan

Timestamp: Wed May 02 10:33:27 EDT 2007

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Application No: 10579290

Version No: 1.0

Input Set:

Output Set:

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Finished: 2007-04-25 14:21:40.978  
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Total Warnings: 0  
Total Errors: 0  
No. of SeqIDs Defined: 4  
Actual SeqID Count: 4

ErrCode	Error Description
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# Sequence Listing

<110> OncoMab GmbH  
VOLLMER, Heinz  
MULLER-HERMELINK, Hans-Konrad

<120> Human monoclonal antibody

<130> 043043-0359295

<140> 10579290

<141> 2007-04-25

<150> 10/579,290

<151> 2004-11-12

<150> EP - 03 026 161.4

<151> 2003-11-14

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<170> PatentIn version 3.3

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Gln	Thr	Ala	Ser	Ile	Thr	Cys	Ser	Gly	Asp	Lys	Leu	Gly	Asp	Lys
				20					25					30

Tyr	Ala	Cys	Trp	Tyr	Gln	Gln	Lys	Pro	Gly	Gln	Ser	Pro	Val	Leu
				35					40					45

Val	Ile	Tyr	Gln	Asp	Ser	Lys	Arg	Pro	Ser	Gly	Ile	Pro	Glu	Arg
				50						55				60

Phe Ser Gly Ser Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser  
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Gly Thr Gln Ala Met Asp Glu Ala Asp Tyr Tyr Cys Gln Ala Trp  
80 85 90

Asp Ser Ser Ile Val Val  
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Ser Tyr Val Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly  
1 5 10 15

cag aca gcc agc atc acc tgc tct gga gat aaa ttg ggg gat aaa 90  
Gln Thr Ala Ser Ile Thr Cys Ser Gly Asp Lys Leu Gly Asp Lys  
20 25 30

tat gct tgc tgg tat cag cag aag cca ggc cag tcc cct gtg ctg 135  
Tyr Ala Cys Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Val Leu  
35 40 45

gtc atc tat caa gat agc aag cgg ccc tca ggg atc cct gag cga 180  
Val Ile Tyr Gln Asp Ser Lys Arg Pro Ser Gly Ile Pro Glu Arg  
50 55 60

ttc tct ggc tcc aac tct ggg aac aca gcc act ctg acc atc agc 225  
Phe Ser Gly Ser Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser  
65 70 75

ggg acc cag gct atg gat gag gct gac tat tac tgt cag gcg tgg 270  
Gly Thr Gln Ala Met Asp Glu Ala Asp Tyr Tyr Cys Gln Ala Trp  
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Arg Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser  
20 25 30

Ser Tyr Ala Met His Trp Val Arg Glu Ala Pro Gly Lys Gly Leu  
35 40 45

Glu Trp Val Ala Val Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr  
50 55 60

Ala Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser  
65 70 75

Lys Asn Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp  
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Thr Ala Val Tyr Tyr Cys Ala Arg Asp Arg Leu Ala Val Ala Gly  
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Lys Thr Phe Asp Tyr  
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